205

Figure 7

a.

```
leutrokine-
lipha M DDSTEREQSRLTSCLKKREEMKLKECVSILPRKESPSVRS
                                                                                    Delete portion of line in brackets
                                 Transmembrane
                                                  Region
      SKDGKLLAATLLLALLSCCLTVVSFYQVAALQGDLASLRAE
      LQGHHAEKLPAGAGAPKAGLEEAPAVTAGLKIFEPPAPGEG 123
      164
                                                                                                  134
                                                                                                   <del>77</del> <u>76</u>
                                                                                                        "G" at position
      FNPWELS----FKRGSALEEKENKINKETGYFELVEENNWOPA----ERUGRGLQAQGYGVRIQDAGVVLLLEGGLACLULGGGLACLULGGGAALLANGVEERD--NQLVWESEGLVLGGGAALLANGVEERD--NGLVWETSGIVMV
                                                                                                   200
                                                                                                        197 should be
                                                                                                  ·170
                                                                                                        black text on
                                                                                                   139
                                                                                                        white
                                                                                                        background
                                                                                  E
      201
                                                                                                   176
      E - - T L D - - - - - N N S Q S A G I A K I E E D E I Q L A T PR E N A 268 S H P D R A - - - - - N N S Q S S A G I A K I E E D E I Q L A T PR E N A 234 C Q R E T D E G A E A K E N Y E P I Y L G G V T O T D K G D R E A E B N R P D Y 217 Y P - - - - - G L Q E P Y L H G M Y H G A A T Q I T D T D T D T T T T D G I P H 190
      QISEDGDV TEAL LET L
KLNASPHGTELEFVKT
IDFAESGQVYTEIIAE
IVLS-BSTV TEGAFAT
                                                                                                   285
                                                                                                   250
                                                                                                   233
```

b.

